PUBLIC TRANSLATION INTO ENGLISH
Descriptive Memory
Of the Invention Patent
On
Genetic constructions, cloning methods of the gene encoding for
Recombinant Human Erythropoietin, selection of producing cell lines
and cell cultures for the mass production of Recumbent Human
Erythropoietin
Aplied by:
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For the term of 20 years
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I. Technical description of the invention
The present invention relates to gene constructions, cloning methods
of the gene encoding for Recombinant Human Erythropoietin,
selection of producing cell lines and cell cultures for the mass
production of Recombinant Human Erythropoietin
II. Technical field of the invention
Genomic DNA that leads to EPO production has been reduced in the
present invention to its minimum expression, including only encoding
sequences and the intros comprised by said sequences plus a few
bases at 3' from the stop codon, and excluding any homologous
sequence, either regulatory or not, at 5' from the initial ATG. The prior
art states that the sequences hereby excluded are very important in
order to obtain a better protein productivity
The vectors used in this invention, in contrast to those described to
date for EPO mass production, have extremely simple expression

control genetic elements, and allow a double selection of the
recombinant cells with antibiotics and methotrexate
These elementary genetic constructions unexpectedly make the cells
transfected with them, produce surprisingly high quantities of EPO per
ml of culture medium per day, several times superior to those
described in former patents and publications
Novelty of the invention – Purpose of the patent
The genetic constructions described to present for EPO production
invariably use homologous elements located at 5' from the first
translated ATG codon, explaining in many cases that this presence of
multiple control elements markedly facilitates the protein expression.
The genomic DNA used in the present invention patent to produce
EPO was isolated so as not to use any homologous element located
at 5' from the firs translated ATG codon, so as to allow the expression
control genetic elements located in the plasmids used, operate in a
similar way as they naturally do, that is to say, operating directly on a
codifying gene, without foreign sequences acting between this gene
and the very control sequences
In this way, the combination of control elements and the gene
encoding for EPO, purpose of this patent, operate with high efficiency,
thus achieving high EPO expressions, comparable to and even higher
than those reported by using much more complex and difficult to
manipulate genetic constructions
Additionally, cotransfection with the two vectors described herein,
which confer different resistances, facilitates the selection, genetic
amplification and maintenance of the cotransfected producing cells
Example 1
To obtain genetic constructions the following procedure was

performed:
Preparation of Human Genomic DNA
10 ml of blood in 10 mM EDTA (pH 8) were extracted from a clinically
healthy adult male subject. The blood was transferred in 5 ml aliquots
to two 50 ml tubes, to which 45 ml of a solution containing 0.3 M of
saccharose, 10 mM Tris-HCL (pH 7.5), 5 mM Mg \mbox{Cl}_2 and 1% of Triton
X 100 was added. The resulting solution was stored at 4° C
The solution was then placed on ice for 10 minutes and centrifuged for
10 minutes at 1000 g and at 4°C. The supernatant was discarded and
the pellet was washed up several times with a 0.075 M NaCl solution
containing 0.025 M EDTA (pH 8), followed by centrifugation for 10
minutes at 1000 g and at 4°C
The resulting pellet thus obtained was resuspended in 3 ml of a 10mM
Tris-HCl (pH 8), 400 mM NaCl, 2 mM EDTA (pH 8) solution. 200 μ l of
10 % SDS (sodium dodecyl sulphate) and 500 μ l K proteinase (1
mg/ml in 1 % SDS and 2 mM EDTA pH 8) were then added, and the
solution was incubated overnight at 37°C. After this incubation, 1 ml of
NaCl saturated solution was added; the solution was shaken and then
centrifuged at 2500 g for 15 minutes
The supernatant was transferred to a 15 ml tube where the volume
was duplicated by the addition of isopropanol. The supernatant and
isopropanol were gently mixed by inversion of the tube and stored at
room temperature until a DNA precipitate was formed, which was
"fished" with a Pasteur blent glass pipette
The DNA was placed in a 2 mL tube, and 1 mL of 70 $\%$ ethanol was
added. After the solution was left stand for one minute, the
supernatant was discarded and the precipitate was left to dry. After
drying, the precipitate was suspended in 500 μl of TE (10 mM Tris-

HCl pH 8 - 1 mM EDTA).-----Concentration of DNA solution was calculated by measuring the absorbance at 260 nm of a 1:1000 dilution of this solution. Each unit of optical density was considered to have 50 µg of genomic DNA. Once the concentration was known, a solution was prepared with 500 ng of genomic DNA per μl in TE.-----B. Preparation of the EPO encoding gene, proper for its expression.-----Gene encoding for EPO was prepared from 500 ng of human genomic DNA obtained in Example 1, adding 400 ng of each of the EPO 1 and EPO 2 primers, in an aqueous 2.5 mM solution containing each deoxynucleotide (dATP, dCTP, dGTP and dTTP), 2.5 units of Taq DNA polymerase (Perkin Elmer) in a final volume of 100 µl using the buffer recommended by the manufacturer. A Perkin Elmer-Cetus Thermal Cycler 480 was used and was programmed for 30 cycles of: 1 minute at 93□C, 1 minute at 55□C and 3 minutes at 72□C. From this reaction, a fragment of approximately 2170 bases containing the EPO gene was obtained.-----The sequence of olygonucleotides employed was:-----EPO 1: 5' GAATTCTCGAGATGGGGGTGCACGGTGAG 3' (SEQ No:2), corresponding to the first bases translated from the EPO gene with the addition of a recognition site for the Xho I enzyme and one for the recognition of Eco RI enzyme to the 5' end. These sites were used in the subsequent cloning steps.-----EPO 2: 5' AAGCTTGGACACACCTGGTCATCTG 3' (SEQ No.:3), complementary to the last translated bases and to some of the nonencoding 3' of the EPO gene. A site for the recognition of the Hind III enzyme was added to the 3' end. This site was used in subsequent cloning steps.-----

The sequence obtained is as follows (SEQ No:4):----gaattctcgagatqgggtgcacggtgagtactcgcgggctgggcgctcccgccgcccgggtc cctgtttgagcggggatttagcgcccggctattggccaggaggtggctgggttcaaggaccggc cttggggagtccttggggatggcaaaaacctgacctgtgaaggggacacagtttgggggttga ggggaagaaggtttgggggttctgctgtgccagtggagaggaagctgataagctgataacctgg gcgctggagccaccacttatctgccagaggggaagcctctgtcacaccaggattgaagtttggc cctggctatctgttctagaatgtcctgcctggctgtggcttctcctgtccctgctgtcgctccctctgggc ctcccagtcctgggcgccccaccacgcctcatctgtgacagccgagtcctggagaggtacctctt ggaggccaaggaggccgagaatatcacggtgagaccccttccccagcacattccacagaact tgggaagctagacactgccccctacataagaataagtctggtggccccaaaccatacctgga aactaggcaaggagcaaagccagcagatcctacggcctgtgggccagggccagagccttca gggaccettgactccccgggctgtgtgcatttcagacgggctgtgctgaacactgcagcttgaatg agaatatcactgtcccagacaccaaagttaatttctatgcctggaagaggatggaggtgagttcct gggaaaggtaaaatggagcagcagagatgaggctgcctgggcgcagaggctcacgtctataa tcccaggctgagatggccgagatgggagaattgcttgagccctggagtttcagaccaacctagg gtggtagtcccagatatttggaaggctgaggcgggaggatcgcttgagcccaggaatttgaggct gcagtgagctgtgatcacaccactgcactccagcctcagtgacagagtgaggccctgtctcaaa cactcactcattcattcattcattcaacaagtcttattgcataccttctgtttgctcagcttggtgctt ggggctgctgaggggcaggagggagagggtgacatgggtcagctgactcccagagtccactc

cctgtaggtcggcagcaggccgtagaagtctggcagggcctggccctgctgtcggaagctgtc ctgcggggccaggccctgttggtcaactcttcccagccgtgggagcccctgcagctgcatgtgga taaagccgtcagtggccttcgcagcctcaccactctgcttcgggctctgggagcccaggtgagta gqagcqqacacttctgcttgccctttctgtaagaaggggagaagggtcttgctaaggagtacagg ccatctcccctccagatgcggcctcagctgctccactccgaacaatcactgctgacactttccgca aactetteegagtetaeteeaattteeteeggggaaagetgaagetgtaeaeaggggaggeetge aggacaggggacagatgaccaggtgtgtccaagctt -----The first translated atg codon, as well as the tga "stop" codon are underlined. ------The sequences of restriction sites.-----

C. Cloning and sequencing.-----

A fragment of approximately 2170 base pairs corresponding to the EPO gene was purified and the ends were blunted by treatment with the RNA polymerase Klenow fragment and cloned in the Sma I site of a M13mp18 vector, following standard techniques applied in molecular biology. The recombinant plasmids obtained were cut with the Xho I and Hind III enzymes; the presence of the insert was verified by electrophoresis of the product resulting from the restriction fragments in a 0.8 % agarose gel developed with ethydium bromide stain. A positive clone (two bands, one having approximately 2200 base pairs and the other one corresponding to the linear vector) was chosen and manually sequenced according to the Sanger's technique using a "T7" sequencing kit" (Pharmacia) and with the aid of an automatic sequences Model 370 A Applied Biosystems International. For each sequencing system the protocols recommended by the manufacturers were followed.-----

D. pVex 1 and pDHFR Vectors
1. Construction of pVex 1 Vector
The pVex1 vector was built following the conventional techniques used
in molecular biology. It consisted of:
a. Fragments of the bacteria1 pBR322 vector, which conferred a
bacterial replication origin and resistance to ampicillin, for amplification
and selection of the vector in E. coli
b. Immediately close to 3' of a) an early promoter of the SV40 virus is
located, which allows the expression of the genes cloned at 5' from
this element
c. Immediately close to 3' of b) the Xho I and Hind III cloning sites are
located, which allow insertion of the genes to be expressed
d. Immediately close to 3' of c) the polyadenylation signal of the SV40
virus is located, which allows the proper polyadenylation of the specific
transcripts of the gene cloned in c)
e. Immediately close to 3' of d) the TK promoter and the gene coding
for neomycin phosphotransferase plus the polyadenylation signal are
located to allow the selection of stably transfected cells through
selection by resistance to neomycin and neomycin-derived antibiotics
such as genetycin. The 3' end of e) is linked to the 5' end of a)
2. pDHFR Vector
The pDHFR vector confers resistance to ampicillin to aid in selection
in bacteria. and includes the DNA copy encoding for mice
dehydrofolate reductase (DHFR), whose expression is controlled by
the SV40 virus early promoter and its polyadenylation signal
The coexpression of DHFR and the EPO-encoding gene allows,
through selection by adding methotrexate (MTX) to the culture
medium, several times-amplification of EPO expression achieved with

the pVex 1-EPO vector
E. Cloning of the EPO-Encoding gene into an Expression Vector-
The sequenced gene was removed by cleavage with the Xho I-Hind III
enzymes of the vector where it was cloned in Example 3. It was then
isolated and cloned in the same restriction sites of the pVex I vector
A positive pVex-EPO clone was isolated. All these operations were
performed according to the conventional genetic engineering
techniques
F. Co-transfection and amplification with MTX
A CHO (Chinese Hamster Ovary) cell line, mutated to be deficient in
the DHFR-enzyme gene (CHO-DHFR), was used to facilitate the
genetic amplification with MTX
During this whole process, cells were grown at 37□ C in a 5% of C0 ₂
atmosphere
CHO cells were cotransfected following the calcium phosphate
technique which, for a 90 mm-diameter Petri dish, consists in:
(a) Replacing the culture medium (alpha-MEM, with10 % of bovine
fetal serum) with fresh medium 4-8 hours before transfection
(b) Adding to a 5 mL tube a 10 g/l HEPES solution (pH 7.1), 16 g/l
NaCl and 10 μ l of a 35 mM Na ₂ HPO ₄ and 35 mM of NaH ₂ PO ₄
solution
(c) Preparing in a separate 1.5 ml tube a solution with 60 μl of 2 M
CaCl ₂ and 10 µg of each DNA to be transfected (pVex-EPO and
pDHFR). Water was added until the volume reached 500 μl. The
pDHFR plasmid described in Example b is based on the pBR 322
plasmid, which confers resistance to ampicillin, can replicate in E.coli,
has the DHFR gene cloned between the early promoter and the
terminator of the SV40, and allows the expression of the DHFR protein

fetal serum. It was essential to perform the dialysis process according to the following schedule: for 100 ml of serum, the serum is placed in a dialysis bag with a porosity under 3000 Da (with a higher porosity, growth factors would be lost, and the cells would not be able to grow and reproduce), the bag is hermetically closed, and completely immersed in a recipient with 5 liters of bidistilled water; where it is left at 4°C for 12 hours. After this, the water is discarded and again 5 liters are added and the bag is then left stand at 4°C for an additional 12 hour period. Then the dialysis bag is removed and the serum is recovered. Dialysis during shorter periods or with smaller volumes, or without replacing the water, would be worthless since a small amount of nucleotides could be left in serum, and therefore the selection with MTX would not work. Dialysis during longer periods would also be worthless since some proteins, necessary for cell growth, could precipitate and be lost.-----G. Isolation of mass production cell lines-----

In order to identify the protein obtained, it was proceeded as described

	in I
	H. Verification of the Specific messenger RNA Sequence
	Produced by the Recombinant Cells
	1 Preparation of RNA from cells
7	otal RNA was prepared from producing cell lines, according to the
1	following protocol:
-	90-mm diameter Petri dish with confluent cells was washed twice
١	with 10 ml of PBS
	- Two ml of GTC buffer were then added and spread all over the dish.
	The GTC buffer was composed of: (1) 50 g guanidinum thiocyanate;
	(2) 0.5 g N-Lauroilsarcosin; (3) 2.5 ml 1 M sodium citrate, pH 7; (4) 0.7
	ml β-mercapthoethanol; (5) 0.33 ml 30% antifoam agent (SIGMA); (6)
I	H ₂ O q.s 100 ml, pH 7.0
	Cells were lysed resulting in a highly viscous solution. The solution
	was transferred to a 15 ml tube, and the process above described was
١	repeated once more using 2 ml of GTC buffer
	- The 15 ml tube was vigorously shaken for 1 minute to break the
	DNA. Fractioning in a cesium chloride gradient was then performed
	For that purpose, 4 ml of a solution containing CsCl (95.97 g CsCl
ć	and 2.5 ml of 1 M Sodium Acetate, pH 5.4, water was added to reach
	a volume of 100 ml) were poured in an ultracentrifuge tube. Over this
	solution and without mixing, the suspension of the cells in GTC was
	added. The tube was then filled with GTC buffer and ultracentrifuged
	at 20°C, for 20 hours at 31000 rpm
	- The RNA remained at the bottom of the tube (pellet) and the DNA
	formed a band in the middle of the cesium chloride gradient
	- The supernatant was discarded, taking special care to eliminate all of
	the DNA. The RNA-containing pellet was left to dry for 5 minutes

,	- The pellet was dissolved in 200 ml of water and transferred to a 1.5
	ml tube
	- 200 ml of 0.4 M Sodium Acetate, pH 4.8, and 2 volumes of ethanol
	were then added, the resulting solution was thoroughly mixed and left to stand for 30 minutes at -80°C.
	- The solution was then centrifuged in a microcentrifuge at 14000 rpm
	for 15 minutes, the supernatant was discarded and the precipitate was
	rinsed with 1 ml of 80 % ethanol
	- The pellet was dried and redissolved in 100 ml of water
	- The concentration of a 1:100 dilution of the RNA solution was
	measured at 260 nm (one optical density unit is equivalent to 40 mg of RNA).
	•
	NOTE: All the solutions and elements used were RNAase-free
	2. Preparation of specific cDNA
	Specific cDNA was prepared following the directions of a kit intended
	for that purpose (cDNA Synthesis System Plus, Amersham - cat. RPN
	1256). EPO2 oligonucleotide was used as specific the primer
	3. Cloning of cDNA Encoding for EPO
	1/20th of the obtained cDNA was amplified using 400 ng of each the
	EPO2 and EPO3 oligonucleotides, and 2.5 mM of each
	deoxynucleotide in the proper buffer, and 2.5 units of Taq DNA
	polymerase, in a total volume of 100 ml
	Thirty five amplification cycles were performed as follows: 1 minute at
	93°C, 1 minute at 55°C and 1 minute at 72°C
	EPO 3 was synthesized as described for EPO 1 and EPO-2, and its
	sequence (5' GAATTCCATGGGGGTGCACGAATGTCC 3') (SEQ ID
	NO:5) corresponded to the first 20 bases encoding for the EPO cDNA,
	adding one site for the recognition of the Eco RI enzyme, to facilitate

	the subsequent genetic manipulations
	A fragment of approximately 600 base pairs was obtained, which was
	cloned in M13mp18 and M13mp19 vectors
	The presence of the insert in the clones with restriction fragments was
	assayed and sequenced in both directions to obtain the complete
	sequence, using the Sanger's method
	Due to the very high autocomplementarity of some regions of the
	gene, which gives rise to many and very ambiguous compressions in
	the radioautography, a sequencing kit using Taq DNA polymerase and
	modified bases was used. Lower quality results were obtained, but the
	compressions were eliminated. The kit used was the Pharmacia-LKB
	Biotechnology Gene aTaq
	The complete sequence of the human erythropoietin DNA copy,
	isolated and cloned, showed to encode for EPO. Therefore, no
	mistakes in the gene or in its transcription were possible
	I. Study of the EPO Produced
	The EPO obtained by culturing the cells of this example underwent
	different quality and identification assays
	1. In a denaturing SDS-PAGE gel the EPO was identified as a wide
	band of molecular weight superior to 30 kDa. See Figure 1
	2. The band was recognized by a monoclonal antibody and by a
	polyclonal antibody to human EPO in a "Western blot" assay. See
	Figure 2
	3.Treatment with glycanases proved the existence of the glycosidio
•	chains whose quantity and molecular weight were as expected. See
	Figure 3
	4. The EPO produced proved to be composed of a series of species
	with isoelectric points ranging from 3.0 to 4.5

5.The complete amino acid sequence of the isolated protein, purified from the culture supernatant of transfected cell lines showed total homology with natural human erythropoietin whose 165 aminoacid sequence is as follows (SEQ No:1):------

sequence is as follows (SEQ No:1):										
NH ₂	Ala	Pro	Pro	Arg	Leu	lle	Cys	Asp	Ser	Arg
	Val	Leu	Glu	Arg	Tyr	Leu	Leu	Glu	Ala	Lys
	Glu	Ala	Glu	<u>Asn</u>	lle	Thr	Thr	Gly	Cys	Ala
	Glu	Hys	Cys	Ser	Leu	Asn	Glu	<u>Asn</u>	lle	Thr
	Val	Pro	Asp	Thr	Lys	Val	Asn	Phe	Tyr	Ala
	Trp	Lys	Arg	Met	Glu	Val	Gly	Gln	Gln	Ala
	Val	Glu	Val	Trp	Gln	Gly	Leu	Ala	Leu	Leu
	Ser	Glu	Ala	Val	Leu	Arg	Gly	Gln	Ala	Leu
	Leu	Val	<u>Asn</u>	Ser	Ser	Gln	Pro	Trp	Glu	Pro
	Leu	Gln	Leu	Hys	Val	Asp	Lys	Ala	Val	Ser
	Gly	Leu	Arg	Ser	Leu	Thr	Thr	Leu	Leu	Arg
	Ala	Leu	Gly	Ala	Gln	Lys	Glu	Ala	lle	Ser
	Pro	Pro	Asp	Ala	Ala	<u>Ser</u>	Ala	Ala	Pro	Leu
	Arg	Thr	lle	Thr	Ala	Asp	Thr	Phe	Arg	Lys
	Gly	Lys	Leu	Lys	Leu	Tyr	Thr	Gly	Glu	Ala

Thr Gly Arg---- COOH .----Sys Arg Asp X Glycosilation sites.----6. The presence of the four glycosilation sites on the 165 amino acid chain, as well as the complex carbohydrate structure, specifically, the sialic acid terminal residues, were demonstrated, as well as its correct in vivo biological activity, when assayed by the model of the ex-hypoxic polycythemic mouse assay, which showed a total parallelism versus the corresponding international standard.-----Productivity achieved, measured by a specific immunoassay, was 50 mg per liter of culture per day.-----What Is Claimed Is:-----Having described and exemplified the nature and main subject of this invention, as well as the manner in which it can be operated, it is hereby stated to claim as of exclusive property and rights:-----1) GENETIC CONSTRUCTIONS, CLONING METHODS OF THE GENE ENCODING FOR RECOMBINANT HUMAN ERYTHROPOIETIN, SELECTION OF PRODUCING CELL **CULTURES** FOR AND CELL THE MASS LINES OF RECOMBINANT HUMAN **PRODUCTION** ERYTHROPOEITIN. characterized by the following: a) the genetic constructions use only the EPO human gene encoding region, without including homologous genenetic elements located at 5' from the first translated ATG codon; b) genetic constructions have viral promoters and terminators, as expression control systems; c) the cloning method for the codifing gene uses directly genomic DNA; d) the cell lines are

selected using a double system, I) resistance to genetycin and

II) resistance to increasing quantities of Methotrexate and e) selected cells EPO productivity is larger than 50 mg/liter of culture medium/day.-----The sequence used is as follows:----gaattctcgagatggggtgcacggtgagtactcgcgggctgggcgctcccgccgc ccgggtccctgtttgagcggggatttagcgcccggctattggccaggaggtggctgg gcctccacgtgccagcggggacttggggagtccttggggatggcaaaaacctgac ctgtgaaggggacacagtttgggggttgaggggaagaaggtttgggggttctgctgtg ccagtggagaggaagctgataagctgataacctgggcgctggagccaccacttatct gccagaggggaagcctctgtcacaccaggattgaagtttggccggagaagtggatg ctctcagcctggctatctgttctagaatgtcctgcctggctgtggcttctcctgtccctgctg tcgctccctctgggcctcccagtcctgggcgccccaccacgcctcatctgtgacagcc gagtcctggagaggtacctcttggaggccaaggaggccgagaatatcacggtgag accccttccccagcacattccacagaactcacgctcagggcttcagggaactcctcc cagatccaggaacctggcacttggtttggggtggagttgggaagctagacactgccc ccctacataagaataagtctggtggccccaaaccatacctggaaactaggcaagga gcaaagccagcagatcctacggcctgtgggccagggccagagccttcagggaccc ttgactccccgggctgtgtgcatttcagacgggctgtgctgaacactgcagcttgaatg agaatatcactgtcccagacaccaaagttaatttctatgcctggaagaggatggaggt gagttcctttttttttttttttttttttggagaatctcatttgcgagcctgattttggatgaaa gggagaatgatcgggggaaaggtaaaatggagcagcagagatgaggctgcctgg gcgcagaggctcacgtctataatcccaggctgagatggccgagatgggagaattgc ttgagccctggagtttcagaccaacctaggcagcatagtgagatcccccatctctaca aacatttaaaaaaattagtcaggtgaagtggtgcatggtggtagtcccagatatttgga

aggctgaggcgggaggatcgcttgagcccaggaatttgaggctgcagtgagctgtg atcacaccactgcactccagcctcagtgacagagtgaggccctgtctcaaaaaaga ctcactcactcattcattcattcattcaacaagtcttattgcataccttctgtttgctca gcttggtgcttggggctgctgaggggcaggaggggagagggtgacatgggtcagctg actcccagagtccactccctgtaggtcgggcagcaggccgtagaagtctggcaggg cctggccctgctgtcggaagctgtcctgcggggccaggccctgttggtcaactcttccc agccgtgggagcccctgcagctgcatgtggataaagccgtcagtggccttcgcagc ctcaccactctgcttcgggctctgggagcccaggtgagtaggagcggacacttctgct tgccctttctgtaagaaggggagaagggtcttgctaaggagtacaggaactgtccgta ctccctccagatgcggcctcagctgctccactccgaacaatcactgctgacactttcc gcaaactcttccgagtctactccaatttcctccggggaaagctgaagctgtacacagg ggaggcctgcaggacaggggacagatgaccaggtgtgtccaagctt -----The first translated atg codon, as well as the tga "stop" codon are underlined. The sequences of restriction sites utilized in the cloning are shown in bold italics.-----CONSTRUCTIONS, according to claim GENETIC characterized because the expression genetic systems consist in two vectors that have as control elements of the EPO expression only the early promoter of the SV40 virus and its terminator, and that allow to produce surprisingly high amounts of EPO, higher than 50/mg/ml/day, in CHO cells stably cotransfected with these vectors and selected according to their resistance to increasing quantities of MTX.-----CONSTRUCTIONS, according to claim 1, GENETIC 3. characterized because the EPO thus obtained consists in 165 amino acids according to the following sequence:-----

NH₂--- Ala

Pro

Pro

Arg

Leu

Ile

Cys

Ser	Arg	Val	Leu	Glu	Arg	Tyr	Leu
Leu	Glu	Ala	Lys	Glu	Ala	Glu	<u>Asn</u>
Ile	Thr	Thr	Gly	Cys	Ala	Glu	Hys
Cys	Ser	Leu	Asn	Glu	<u>Asn</u>	Ile	Thr
Val	Pro	Asp	Thr	Lys	Val	Asn	Phe
Tyr	Ala	Trp	Lys	Arg	Met	Glu	Val
Gly	Gln	Gln	Ala	Val	Glu	Val	Trp
Gln	Gly	Leu	Ala	Leu	Leu	Ser	Glu
Ala	Val	Leu	Arg	Gly	Gln	Ala	Leu
Leu	Val	<u>Asn</u>	Ser	Ser	Gln	Pro	Trp
Glu	Pro	Leu	Gln	Leu	Hys	Val	Asp
Lys	Ala	Val	Ser	Gly	Leu	Arg	Ser
Leu	Thr	Thr	Leu	Leu	Arg	Ala	Leu
Gly	Ala	Gln	Lys	Glu	Ala	Ile	Ser
Pro	Pro	Asp	Ala	Ala	<u>Ser</u>	Ala	Ala
Pro	Leu	Arg	Thr	Ile	Thr	Ala	Asp
Thr	Phe	Arg	Lys	Leu	Phe	Arg	Val
Tyr	Ser	Asn	Phe	Leu	Arg	Gly	Lys
Leu	Lys	Leu	Tyr	Thr	Gly	Glu	Ala
Cys	Arg	Thr	Gly	Asp	СООН	,	

----- X: glycosilation sites.----

There follows an illegible signature followed by a seal that reads: HUMBERTO M. DE PASQUALE. ATTORNEY.-----

IX. Abstract.-----

The gene encoding for human erythropoietin (EPO) was obtained from human genomic DNA. This gene does not include homologous genetic elements located at 5' from the first translated ATG codon. The gene was cloned in an expression plasmid for eukaryotic cells. Hamster Ovary Cells (CHO) were cotransfected with this vector and other plasmid

expressing for dehydrofolatereductase (DHFR). Both plasmids have as unique expression control elements the early promoter of the SV40 virus and its polyadenylation signal. This system allows a double selection of the recombinant cells, as it confers resistance to geniticine and to methotrexate (MTX). Once selection was made, genetic amplification with MTX was performed. Productive capacity of the selected recombinant cells, was tested by means of an immunoassay specific for EPO-----TECHNICAL DATASHEET-----(10) PUBLICATION No.: AR------There appears a seal that reads I.N.P.I. 6 Nov 1998 12 37.---Reception Desk.-----(21) APPLICATION No.:-----On the left, there appears the Argentinean Coat of Arms -----(19) ------(51)INT. CL:----I.N.P.I.-----Argentine Republic-----P98 0105609.-----(12) X INVENTION PATENT------UTILITY MODEL-----(2) SUBMISSION DATE: -----(30) PRIORITY DATA:-----(41) APPLICATION PUBLICATION DATE: -----BULLETIN No.:-----(61) ADDITIONAL TO: ------(62) DIVISIONAL FROM:-----(71) APPLICANT: BIO SIDUS S.A.-----

(72) Inventor(s):
(74) Agent. 611
(83) Microorganism deposit:
(54) TITLE OF THE INVENTION: "GENETIC
CONSTRUCTIONS, CLONING METHOD FOR THE
ENCODING GENE FOR RECOMBINANT HUMAN
ERYTHROPOIETIN, SELECTION OF PRODUCING CELL
LINE AND CELL CULTURE FOR THE MASS PRODUCTION
OF RECOMBINANT HUMAN ERYTROPOIETIN."
(57) ABSTRACT:
The encoding gene for recombinant erythropoietin (EPO) was
obtained from human genomic DNA. This gene does not
include homologous genetic elements located at the 5' of the
first ATG translated codon. It was cloned in an expression
vector for eukaryotic cells. Chinese Hamster Ovary (CHO) cell
were co-transfected with this vector and other expression
plasmid for dehydrofolactoreductase (DHFR). Both plasmids
have the early promoter of SV40 virus and its polyadenylation
signal. This system allows a double selection of recombinant
cells, since they confer a geniticine and methotrexate (MTX)
resistance. Once the selection is performed, gene
amplification was done with MTX. Testing of the producing
capacity of the chosen recombinant cells was performed with
a specific immunoassay for EPO
Argentine Republic. There appears an Argentinean Coat of
Arms. I.N.P.I
APPLICATION OF
INVENTION PATENT: X

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UTILITY MODEL CERTIFICATE
There appears a seal that reads INPI. 6 Nov. 1998. Reception
Desk
FILING DATE
Proceedings No
I. Applicant
(1) Name and Surname / Company's Name:
BIO SIDUS S.A
2). Identity Document:
Marital Status:
Marriage:
Spouse's Name:
3) Retirement Account:
No of CUIL (Unique Labor Identification Code) or CUIT
(Unique Tax Identification Code): 30-59811709-4 V.A.T
Liable for V.A.T Registration
4) Registered in the National Industrial Registry (Decree-Law
19.971/72) NoDoes not apply
5) Real Address: Constitución 4234. Buenos Aires. Argentina.
Legal Address: Alsina 971 – 1º Piso, of. "10, Buenos Aires
II. OBJECT:
6) Title of invention: "Genetic constructions, cloning method
for the encoding gene for recombinant human erythropoietin
selection of producing cell lines and cell culture for the mass
production of recombinant human erythropoietin
7) Type of patent:
a) Final:for a 20-year period
b) Additional to Patent No.:

8) Act 17,011. Priority Date:
Country:
No
III. Attached Documents
(9) It is attached:
a) Fee receipt for the requested service
b) Form in duplicate
c) Cover in duplicate
d) Descriptive Memory in duplicate
e) Signed claims in duplicate
f) 2 copies of the first claim
g) Diagrams in triplicate
h) Number of boards
i) Reduced-scale copies
j) Certified copy (Act 17,011)
k) Assignment Document
I) Draft drawings
IV Legal Entities
10) The corporation, represented by HUMBERTO MARIO DE
PASQUALE
who state under oath that he is the attorney with his powers in
force and that the corporation is registered in the Public
Commerce Registry on Date: 10/07/1983 No. 7258. Page:
Book: 98. Volume: A
V. Power of Attorney
11) Power of Attorney registered inRegistered in
the INPI (National Institute of Industrial Property) under No.:
Other Registry:No

	12) In this case,: CARLOS MIGUEL COLL ARECO and/or
	HUGO EDUARDO MARTINEZ LAHITOU are authorized to
	proceed in this matter until its finalization with the power of
	signing documents, waving, if necessary, and requesting
	certificates
	13) Power of Attorney is attached
	14) Retirement Account / Company: CONSOLIDAR No. CUIL
	or CUIT: CUIT 20-04991729-6 CUIT: 20-16821007-9
	15) Agent No. 611/900
	VI <u>Statement</u>
,	16) In view of the Provision with no number dated 7 June 1901
	(on Patentability in foreign countries) it is declared that the
	invention has not been patented abroad
	VII. Remarks: According to Article 19 Act 24,481 and its
	Regulatory Decree, complementary information will be
	supplied within the legal term
	There appears a signature. Carlos Miguel Coll Areco
	(Signature of the authorized person)
	There appears a signature. Humberto Mario De Pasquale.
	Attorney. Signature of applicant
	There appears the Argentinian Coat of Arms
	MINISTRY OF ECONOMY AND PUBLIC SERVICES
	NATIONAL INSTITUTE OF INDUSTRIAL PROPERTY
	INVENTION PATENT
	February 23 1999
	Reception Desk
	National Institute of Industrial Property
	Application of: Invention PatentFile:

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Under the proceedings P98 01 05609 an INVENTION
PATENT application has been entered
Buenos Aires, (blank)199
INPI 6 Nov. 1998. Reception Desk
Date: 11/06/1999Time: 12:37
Resp: Coll Areco, Carlos MiguelP19990100679
File Bar CodeQuery Priority Order
01996010560906111237P19980105609-
I, VIVIANA BEATRIZ CUMBO, A SWORN PUBLIC
TRANSLATOR, DO HEREBY CERTIFY THE FOREGOING
TO BE A TRUE TRANSLATION INTO ENGLISH OF THE
PHOTOCOPY OF THE ORIGINAL DOCUMENT IN SPANISH
LANGUAGE, ATTACHED HERETO, WHICH I HAVE HAD
BEFORE ME. DONE AND SIGNED IN BUENOS AIRES, ON
THIS FOURTEENTH DAY OF NOVEMBER, TWO
THOUSAND AND THREE
YO, VIVIANA BEATRIZ CUMBO, TRADUCTORA PÚBLICA
MATRICULADA, CERTIFICO POR LA PRESENTE, QUE
ÉSTA ES TRADUCCIÓN FIEL AL IDIOMA INGLÉS DE LA
FOTOCOPIA DEL DOCUMENTO ORIGINAL REDACTADO
EN IDIOMA CASTELLANO, ADJUNTA A LA PRESENTE,
QUE HE TENIDO A LA VISTA, Y A LA CUAL ME REMITO,
EN BUENOS AIRES, A LOS CATORCE DÍAS DEL MES DE
NOVIEMBRE DE DOS MIL TRES